## **HS arch Differential expression HvN**

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k141_1	81201_18	ф ФФ ф	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_181201_18pdf)	-7.69	3.73e-09	1.55
k141_3	72642_30	ф • • • • • • • • • • • • • • • • • • •	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_372642_30pdf)	-7.69	1.92e-06	1.14
k141_3	26599_21_CBS_domain	•	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_326599_21_CBS_do…	-7.49	5.62e-07	5.35
k141_9	3077_13	ф Ф	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_93077_13pdf)	-6.72	1.99e-07	4.15
k141_3	26599_17_Universal_stress_protein	<b>•</b> • • • • • • • • • • • • • • • • • •	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_326599_17_Univers	-6.38	6.43e-07	5.35
k141_3	92539_6_COG0589_Universal_stress_protein_Usp	ф • • • • • • • • • • • • • • • • • • •	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_392539_6_COG058	-6.32	5.21e-07	5.35
k141_1	81201_19_Domain_of_unknown_function_DUF302	•	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_181201_19_Domain	-6.16	6.60e-05	2.74
k141_9	6470_27_CBS_domain	•	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_96470_27_CBS_do	-5.63	6.54e-05	2.74
k141_9	2599_11_COG1522_Transcriptional_regulators	• • • • • • • • • • • • • • • • • • •	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_92599_11_COG152	-5.59	1.83e-03	5.170
k141_7	6157_13_Belongs_to_the_P_IIprotein_family	•	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_76157_13_Belongs	-5.57	1.39e-05	7.220
k141_9	2599_5_helix_turn_helix_ASNC_type	Φ		eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_92599_5_helix_turn	-5.12	2.39e-03	6.210
k141_3	26599_18_COG0589_Universal_stress_protein_Us	0 000-0	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_326599_18_COG05	-5.07	2.26e-04	7.85
k141_4	40873_29_Belongs_to_the_small_heat_shock_pro	0 0 0 0		eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_440873_29_Belongs	-4.42	1.29e-03	4.13
k141_4	47824_6	ф ф <del></del>	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_447824_6pdf)	-4.14	1.65e-06	1.14
k141_9	2599_3_Lrp_AsnC_ligand_binding_domain	· · · · · · · · · · · · · · · · · · ·	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_92599_3_trp_AsnC	-4.02	1.89e-04	7.16
k141_4	40873_33_Stabilizes_TBP_binding_to_an_archae		(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_440873_33_Stabilize	-3.20	1.86e-03	5.170
k141_1	5183_34	· · · · · · · · · · · · · · · · · · ·	(figuresHS_prok_arch_RNAs	eq_analysis_with_DESeq2_long_	names_HvN_alldata/boxplot.k141_15183_34pdf)	-2.61	3.97e-03	9.72
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